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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/667,947A

DATE: 06/10/2003 P6

TIME: 11:18:45

Input Set : A:\07039-298001.txt

Output Set: N:\CRF4\06102003\I667947A.raw

4 <110> APPLICANT: Russell, Stephen James
 5 Cattaneo, Roberto
 6 Peng, Kah-Whye
 7 Schneider, Urs
 8 Murphy, Anthea L.
 10 <120> TITLE OF INVENTION: Therapeutic methods and compositions
 using viruses of the recombinant paramyxoviridae family
 14 <130> FILE REFERENCE: 07039-298001
 16 <140> CURRENT APPLICATION NUMBER: US 09/667,947A
 17 <141> CURRENT FILING DATE: 2000-09-22
 19 <150> PRIOR APPLICATION NUMBER: US 60/155,873
 20 <151> PRIOR FILING DATE: 1999-09-24
 22 <160> NUMBER OF SEQ ID NOS: 49
 24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 4
 28 <212> TYPE: PRT
 29 <213> ORGANISM: Artificial Sequence
 W--> 30 <220> FEATURE:
 31 <223> OTHER INFORMATION: Factor Xa cleavage site
 W--> 32 <400> SEQUENCE: 1
 33 Ile Glu Gly Arg
 34 1
 36 <210> SEQ ID NO: 2
 37 <211> LENGTH: 4
 38 <212> TYPE: PRT
 39 <213> ORGANISM: Artificial Sequence
 W--> 40 <220> FEATURE:
 41 <223> OTHER INFORMATION: Furin cleavage site
 W--> 42 <220> FEATURE:
 43 <221> NAME/KEY: VARIANT
 44 <222> LOCATION: 2
 45 <223> OTHER INFORMATION: Xaa = Any 20 amino acids
 W--> 46 <400> SEQUENCE: 2
 W--> 47 Arg Xaa Lys Arg
 48 1
 50 <210> SEQ ID NO: 3
 51 <211> LENGTH: 6
 52 <212> TYPE: PRT
 53 <213> ORGANISM: Artificial Sequence
 W--> 54 <220> FEATURE:
 55 <223> OTHER INFORMATION: MMP cleavage site
 58 <400> SEQUENCE: 3

RAW SEQUENCE LISTING

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59 Pro Leu Gly Leu Trp Ala

60 1 5

62 <210> SEQ ID NO: 4

63 <211> LENGTH: 6

64 <212> TYPE: PRT

65 <213> ORGANISM: Artificial Sequence

W--> 66 <220> FEATURE:

67 <223> OTHER INFORMATION: Caspase-1 cleavage site

W--> 68 <400> SEQUENCE: 4

69 Tyr Glu Val Asp Gly Trp

70 1 5

72 <210> SEQ ID NO: 5

73 <211> LENGTH: 7

74 <212> TYPE: PRT

75 <213> ORGANISM: Artificial Sequence

W--> 76 <220> FEATURE:

77 <223> OTHER INFORMATION: Caspase-2 cleavage site

W--> 78 <400> SEQUENCE: 5

79 Val Asp Val Ala Asp Gly Trp

80 1 5

82 <210> SEQ ID NO: 6

83 <211> LENGTH: 7

84 <212> TYPE: PRT

85 <213> ORGANISM: Artificial Sequence

W--> 86 <220> FEATURE:

87 <223> OTHER INFORMATION: Caspase-3 cleavage site

W--> 88 <400> SEQUENCE: 6

89 Val Asp Gln Met Asp Gly Trp

90 1 5

92 <210> SEQ ID NO: 7

93 <211> LENGTH: 6

94 <212> TYPE: PRT

95 <213> ORGANISM: Artificial Sequence

W--> 96 <220> FEATURE:

97 <223> OTHER INFORMATION: Caspase-4 cleavage site

W--> 98 <400> SEQUENCE: 7

99 Leu Glu Val Asp Gly Trp

100 1 5

102 <210> SEQ ID NO: 8

103 <211> LENGTH: 6

104 <212> TYPE: PRT

105 <213> ORGANISM: Artificial Sequence

W--> 106 <220> FEATURE:

107 <223> OTHER INFORMATION: Caspase-6 cleavage site

W--> 108 <400> SEQUENCE: 8

109 Val Gln Val Asp Gly Trp

110 1 5

112 <210> SEQ ID NO: 9

113 <211> LENGTH: 7

RAW SEQUENCE LISTING
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Input Set : A:\07039-298001.txt
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114 <212> TYPE: PRT
115 <213> ORGANISM: Artificial Sequence
W--> 116 <220> FEATURE:
117 <223> OTHER INFORMATION: Caspase-7 cleavage site
W--> 118 <400> SEQUENCE: 9
119 Val Asp Gln Val Asp Gly Trp
120 1 5
122 <210> SEQ ID NO: 10
123 <211> LENGTH: 4
124 <212> TYPE: PRT
125 <213> ORGANISM: Artificial Sequence
W--> 126 <220> FEATURE:
127 <223> OTHER INFORMATION: Proprotein convertase cleavage site
W--> 128 <400> SEQUENCE: 10
129 Arg Gly Leu Thr
130 1
132 <210> SEQ ID NO: 11
133 <211> LENGTH: 17
134 <212> TYPE: PRT
135 <213> ORGANISM: Artificial Sequence
W--> 136 <220> FEATURE:
137 <223> OTHER INFORMATION: FMDV protease 2A cleavage site
W--> 138 <400> SEQUENCE: 11
139 Asn Phe Asp Leu Leu Lys Leu Ala Gly Asp Val Glu Ser Asn Pro Gly
140 1 5 10 15
141 Pro
143 <210> SEQ ID NO: 12
144 <211> LENGTH: 34
145 <212> TYPE: PRT
146 <213> ORGANISM: Paramyxoviridae
W--> 147 <220> FEATURE:
148 <223> OTHER INFORMATION: H protein cytoplasmic tail
W--> 149 <400> SEQUENCE: 12
150 Met Ser Pro Gln Arg Asp Arg Ile Asn Ala Phe Tyr Lys Asp Asn Pro
151 1 5 10 15
152 His Pro Lys Gly Ser Arg Ile Val Ile Asn Arg Glu His Leu Met Ile
153 20 25 30
154 Asp Arg
157 <210> SEQ ID NO: 13
158 <211> LENGTH: 33
159 <212> TYPE: PRT
160 <213> ORGANISM: Paramyxoviridae
W--> 161 <220> FEATURE:
162 <223> OTHER INFORMATION: F protein cytoplasmic tail
W--> 163 <400> SEQUENCE: 13
164 Arg Gly Arg Cys Asn Lys Lys Gly Glu Gln Val Gly Met Ser Arg Pro
165 1 5 10 15
166 Gly Leu Lys Pro Asp Leu Thr Gly Thr Ser Lys Ser Tyr Val Arg Ser
167 20 25 30

RAW SEQUENCE LISTING
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168 Leu
172 <210> SEQ ID NO: 14
173 <211> LENGTH: 5
174 <212> TYPE: PRT
175 <213> ORGANISM: Artificial Sequence
W--> 176 <220> FEATURE:
177 <223> OTHER INFORMATION: Furin cleavage site
W--> 178 <400> SEQUENCE: 14
179 Arg Arg His Lys Arg
180 1 5
182 <210> SEQ ID NO: 15
183 <211> LENGTH: 4
184 <212> TYPE: PRT
185 <213> ORGANISM: Artificial Sequence
W--> 186 <220> FEATURE:
187 <223> OTHER INFORMATION: Furin cleavage site
W--> 188 <400> SEQUENCE: 15
189 Arg His Lys Arg
190 1
192 <210> SEQ ID NO: 16
193 <211> LENGTH: 47
194 <212> TYPE: DNA
195 <213> ORGANISM: Artificial Sequence
W--> 196 <220> FEATURE:
197 <223> OTHER INFORMATION: Primer
W--> 198 <400> SEQUENCE: 16
199 ttttcctttt gcggccgctt tcataaacgc ttctgcaggg acccctc 47
201 <210> SEQ ID NO: 17
202 <211> LENGTH: 56
203 <212> TYPE: DNA
204 <213> ORGANISM: Artificial Sequence
W--> 205 <220> FEATURE:
206 <223> OTHER INFORMATION: Primer
W--> 207 <400> SEQUENCE: 17
208 gtccatgcgg cccagccggc ccgattaaag agagaggcag aggacctgca ggtggg 56
210 <210> SEQ ID NO: 18
211 <211> LENGTH: 18
212 <212> TYPE: PRT
213 <213> ORGANISM: Artificial Sequence
W--> 214 <220> FEATURE:
215 <223> OTHER INFORMATION: Amino acid sequence coded for by primer
W--> 216 <400> SEQUENCE: 18
217 Val His Ala Ala Gln Pro Ala Arg Leu Lys Arg Glu Ala Glu Asp Leu
218 1 5 10 15
219 Gln Val
222 <210> SEQ ID NO: 19
223 <211> LENGTH: 50
224 <212> TYPE: DNA
225 <213> ORGANISM: Artificial Sequence

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RAW SEQUENCE LISTING
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Input Set : A:\07039-298001.txt
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W--> 226 <220> FEATURE:
 227 <223> OTHER INFORMATION: Primer
 229 <400> SEQUENCE: 19
 230 ttttccttt gcggccgctt tcatcatcaa cgcttctgca gggaccctc 50
 232 <210> SEQ ID NO: 20
 233 <211> LENGTH: 59
 234 <212> TYPE: DNA
 235 <213> ORGANISM: Artificial Sequence
 W--> 236 <220> FEATURE:
 237 <223> OTHER INFORMATION: Primer
 W--> 238 <400> SEQUENCE: 20 59
 239 gtccatgcgg cccagccggc cggtggaggc ggttcagagg cagaggacct gcaggtggg
 241 <210> SEQ ID NO: 21
 242 <211> LENGTH: 19
 243 <212> TYPE: PRT
 244 <213> ORGANISM: Artificial Sequence
 W--> 245 <220> FEATURE:
 246 <223> OTHER INFORMATION: Amino acid sequence coded for by primer
 W--> 247 <400> SEQUENCE: 21
 248 Val His Ala Ala Gln Pro Ala Gly Gly Gly Ser Glu Ala Glu Asp
 249 1 5 10 15
 250 Leu Gln Val
 253 <210> SEQ ID NO: 22
 254 <211> LENGTH: 16
 255 <212> TYPE: PRT
 256 <213> ORGANISM: Paramyxoviridae
 W--> 257 <220> FEATURE:
 258 <223> OTHER INFORMATION: F protein cytoplasmic tail
 W--> 259 <400> SEQUENCE: 22
 260 Arg Gly Arg Cys Asn Lys Lys Gly Glu Gln Gly Met Ser Arg Pro Gly
 261 1 5 10 15
 263 <210> SEQ ID NO: 23
 264 <211> LENGTH: 9
 265 <212> TYPE: PRT
 266 <213> ORGANISM: Paramyxoviridae
 W--> 267 <220> FEATURE:
 268 <223> OTHER INFORMATION: Cytoplasmic tail of Fc(24 mutant
 W--> 269 <400> SEQUENCE: 23
 270 Arg Gly Arg Cys Asn Lys Lys Gly Glu
 271 1 5
 273 <210> SEQ ID NO: 24
 274 <211> LENGTH: 26
 275 <212> TYPE: DNA
 276 <213> ORGANISM: Artificial Sequence
 W--> 277 <220> FEATURE:
 278 <223> OTHER INFORMATION: Primer
 W--> 279 <400> SEQUENCE: 24 26
 280 aaaactgcag actcaaaggt caatgc
 282 <210> SEQ ID NO: 25

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 06/10/2003
PATENT APPLICATION: US/09/667,947A TIME: 11:18:46

Input Set : A:\07039-298001.txt
Output Set: N:\CRF4\06102003\I667947A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 2
Seq#:35; Xaa Pos. 5,6,7
Seq#:36; N Pos. 1,2,3,4,5,6,7,8,9,10,11,12,15,16,17,20

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/667,947A

DATE: 06/10/2003
TIME: 11:18:46

Input Set : A:\07039-298001.txt
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L:30 M:283 W: Missing Blank Line separator, <220> field identifier
L:32 M:283 W: Missing Blank Line separator, <400> field identifier
L:40 M:283 W: Missing Blank Line separator, <220> field identifier
L:42 M:283 W: Missing Blank Line separator, <220> field identifier
L:46 M:283 W: Missing Blank Line separator, <400> field identifier
L:47 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
L:54 M:283 W: Missing Blank Line separator, <220> field identifier
L:66 M:283 W: Missing Blank Line separator, <220> field identifier
L:68 M:283 W: Missing Blank Line separator, <400> field identifier
L:76 M:283 W: Missing Blank Line separator, <220> field identifier
L:78 M:283 W: Missing Blank Line separator, <400> field identifier
L:86 M:283 W: Missing Blank Line separator, <220> field identifier
L:88 M:283 W: Missing Blank Line separator, <400> field identifier
L:96 M:283 W: Missing Blank Line separator, <220> field identifier
L:98 M:283 W: Missing Blank Line separator, <400> field identifier
L:106 M:283 W: Missing Blank Line separator, <220> field identifier
L:108 M:283 W: Missing Blank Line separator, <400> field identifier
L:116 M:283 W: Missing Blank Line separator, <220> field identifier
L:118 M:283 W: Missing Blank Line separator, <400> field identifier
L:126 M:283 W: Missing Blank Line separator, <220> field identifier
L:128 M:283 W: Missing Blank Line separator, <400> field identifier
L:136 M:283 W: Missing Blank Line separator, <220> field identifier
L:138 M:283 W: Missing Blank Line separator, <400> field identifier
L:147 M:283 W: Missing Blank Line separator, <220> field identifier
L:149 M:283 W: Missing Blank Line separator, <400> field identifier
L:161 M:283 W: Missing Blank Line separator, <220> field identifier
L:163 M:283 W: Missing Blank Line separator, <400> field identifier
L:176 M:283 W: Missing Blank Line separator, <220> field identifier
L:178 M:283 W: Missing Blank Line separator, <400> field identifier
L:186 M:283 W: Missing Blank Line separator, <220> field identifier
L:188 M:283 W: Missing Blank Line separator, <400> field identifier
L:196 M:283 W: Missing Blank Line separator, <220> field identifier
L:198 M:283 W: Missing Blank Line separator, <400> field identifier
L:205 M:283 W: Missing Blank Line separator, <220> field identifier
L:207 M:283 W: Missing Blank Line separator, <400> field identifier
L:214 M:283 W: Missing Blank Line separator, <220> field identifier
L:216 M:283 W: Missing Blank Line separator, <400> field identifier
L:226 M:283 W: Missing Blank Line separator, <220> field identifier
L:236 M:283 W: Missing Blank Line separator, <220> field identifier
L:238 M:283 W: Missing Blank Line separator, <400> field identifier
L:245 M:283 W: Missing Blank Line separator, <220> field identifier
L:247 M:283 W: Missing Blank Line separator, <400> field identifier
L:257 M:283 W: Missing Blank Line separator, <220> field identifier
L:259 M:283 W: Missing Blank Line separator, <400> field identifier
L:267 M:283 W: Missing Blank Line separator, <220> field identifier
L:269 M:283 W: Missing Blank Line separator, <400> field identifier
L:277 M:283 W: Missing Blank Line separator, <220> field identifier
L:279 M:283 W: Missing Blank Line separator, <400> field identifier

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PATENT APPLICATION: US/09/667,947A

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Input Set : A:\07039-298001.txt
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L:289 M:283 W: Missing Blank Line separator, <400> field identifier
L:296 M:283 W: Missing Blank Line separator, <220> field identifier
L:298 M:283 W: Missing Blank Line separator, <400> field identifier
L:389 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:35
L:392 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:35
L:393 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:409 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:412 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:415 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:418 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:421 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:424 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:427 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:430 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:433 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:436 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:439 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:442 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:445 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:446 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0